



Entrez PubMed Nucleotide Protein Genome Structure PMC Taxonomy

Search Nucleotide

20  Text

All: 8807 bacteria: 2 mRNA: 4879 RefSeq: 360

Entrez Nucleotide

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[NM\\_101672](#)

Arabidopsis thaliana mother of FT and TF1 protein (MFT) (At1g18100)  
mRNA, complete cds  
gi|42562142|ref|NM\_101672.2|[42562142]

Related resources

☐ 2: [Reports](#) [Links](#)  
[NM\\_105222](#)

Arabidopsis thaliana flowering locus T protein (FT) (At1g65480) mRNA,  
complete cds  
gi|30697201|ref|NM\_105222.2|[30697201]

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[NM\\_120465](#)

Arabidopsis thaliana terminal flower 1 protein (TFL1) (At5g03840) mRNA,  
complete cds  
gi|30680238|ref|NM\_120465.2|[30680238]

☐ 4: [Reports](#) [Links](#)  
[NM\\_120208](#)

Arabidopsis thaliana phosphatidylethanolamine-binding family protein  
(At5g01300) mRNA, complete cds  
gi|30679209|ref|NM\_120208.3|[30679209]

☐ 5: [Reports](#) [Links](#)  
[NM\\_125597](#)

Arabidopsis thaliana brother of FT and TFL1 protein (BFT) (At5g62040)  
mRNA, complete cds  
gi|18424542|ref|NM\_125597.1|[18424542]

☐ 6: [Reports](#) [Links](#)  
[BC089332](#)

Mus musculus phosphatidylethanolamine binding protein, mRNA (cDNA  
clone MGC:102065 IMAGE:30359654), complete cds  
gi|59808980|gb|BC089332.1|[59808980]

- 7:**  
AY730275 Reports Links  
Homo sapiens cousin-of-RKIP 1 protein (CORK1) mRNA, complete cds  
gi|57864776|gb|AY730275.1|[57864776]
- 8:**  
AB188117 Reports Links  
Takifugu rubripes CECR5 mRNA for cat eye syndrome candidate region 5 protein, complete cds  
gi|59857477|dbj|AB188117.1|[59857477]
- 9:**  
AJ785002 Reports Links  
Echinococcus multilocularis raf gene for raf serine/threonine kinase, exons 1-9  
gi|50724570|emb|AJ785002.1|[50724570]
- 10:**  
AJ785001 Reports Links  
Echinococcus multilocularis mRNA for raf serine/threonine kinase (raf gene)  
gi|50724568|emb|AJ785001.1|[50724568]
- 11:**  
AJ785000 Reports Links  
Echinococcus multilocularis mRNA for ras GTPase (ras gene)  
gi|50724566|emb|AJ785000.1|[50724566]
- 12:**  
NM\_205743 Reports Links  
Danio rerio serine/threonine protein kinase ARAF standard form (ARAF), mRNA  
gi|45387896|ref|NM\_205743.1|[45387896]
- 13:**  
BC087096 Reports Links  
Rattus norvegicus mitochondrial ribosomal protein L38 (predicted), mRNA (cDNA clone MGC:94655 IMAGE:7191805), complete cds  
gi|56270359|gb|BC087096.1|[56270359]
- 14:**  
NM\_002165 Reports Links  
Homo sapiens inhibitor of DNA binding 1, dominant negative helix-loop-helix protein (ID1), transcript variant 1, mRNA  
gi|31317298|ref|NM\_002165.2|[31317298]
- 15:**  
NM\_181353 Reports Links  
Homo sapiens inhibitor of DNA binding 1, dominant negative

helix-loop-helix protein (ID1), transcript variant 2, mRNA  
gi|31317296|ref|NM\_181353.1|[31317296]

☐ 16:

NM\_001009369

Reports

Links

Rattus norvegicus mitochondrial ribosomal protein L38 (predicted)  
(Mrpl38\_predicted), mRNA  
gi|57164122|ref|NM\_001009369.1|[57164122]

☐ 17:

NM\_012610

Reports

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Rattus norvegicus nerve growth factor receptor (TNFR superfamily, member  
16) (Ngfr), mRNA  
gi|6981271|ref|NM\_012610.1|[6981271]

☐ 18:

NM\_033360

Reports

Links

Homo sapiens v-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog  
(KRAS), transcript variant a, mRNA  
gi|34485724|ref|NM\_033360.2|[34485724]

☐ 19:

NM\_004985

Reports

Links

Homo sapiens v-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog  
(KRAS), transcript variant b, mRNA  
gi|34485723|ref|NM\_004985.3|[34485723]

☐ 20:

XM\_572136

Reports

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Cryptococcus neoformans var. neoformans JEC21 chromosome 7  
gi|58269959|ref|XM\_572136.1|[58269959]

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